
Sequence Listing was accepted.

If you need help call the Patent Electronic Business Center at (866) 217-9197 (toll free).

Reviewer: markspencer

Timestamp: [year=2008; month=3; day=21; hr=16; min=9; sec=17; ms=929;]

Validated By CRFValidator v 1.0.3

Application No: 10589735 Version No: 1.0

Input Set:

Output Set:

Started: 2008-03-11 13:48:27.804 **Finished:** 2008-03-11 13:48:28.306

Elapsed: 0 hr(s) 0 min(s) 0 sec(s) 502 ms

Total Warnings: 8
Total Errors: 0

No. of SeqIDs Defined: 10

Actual SeqID Count: 10

Error code		or code	Error Description									
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(3)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(4)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(5)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(6)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(7)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(8)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(9)
	W	213	Artificial	or	Unknown	found	in	<213>	in	SEO	ID	(10)

SEQUENCE LISTING

<110>	HAM]	IES, Jim ILTON-BRUCE, LAR, Simon	Monica Anr	ne			
<120>	MUTA	ATION ASSOCI	ATED WITH I	LACUNAR STRO	OKES		
<130>	0641	-0285PUS1					
<140>	1058	39735					
<141>	2008	3-03-11					
<150>	PCT/	'AU2005/0002	218				
<151>	2005	5-02-18					
<150>	AU 2	2004900790					
<151>	2004	1-02-18					
<160>	10						
<170>	Pate	entIn versio	on 3.3				
<210>	1						
<211>	1026	5					
<212> DNA							
<213>	Homo	sapiens					
<400>	1						
aaaggac	caga	tattgcagaa	gagagaaggt	ataactggga	ccaaaagcct	tgagaaggaa	60
agagaca	atgg	agcaaatcat	tcacagtaac	agcagacagc	agagaagaga	cacatggttg	120
tacagaç	ggca	cctcctttgg	gtctttactc	aaatgcccca	ttatcagtga	gaacttctct	180
gactgct	gtt	cttcagcaga	gggtattcct	tatccccttt	cttgctttat	gtgttttctc	240
cataaca	atat	gtgcatatcc	ataacacaca	catgcatcac	ctagagcatt	atatatgcca	300
cagtgac	catg	ttttgctgat	ttctcaattg	actcccccat	tggaatgaac	gtaagcttga	360
ggaagac	gtt	ttgtcctgtt	ctgtagcatc	tagaacagcg	cctggcacat	agtaggtact	420
caataaa	atgc	cagctgcatg	aggaaatgaa	tgagctgtgt	gggggatgta	cttgagtgaa	480
ctctaaa	agtc	agagtggtgt	tgagagaaaa	atgcttgaaa	tccagatgtt	ggaaggtgac	540
acagagt	agt	agcctggtga	gaacagttag	atcttagggg	ttcctactac	agccctccct	600
tccgcac	ctt	tttggctgtc	accatgatca	agctactgaa	tctctctgag	acgcaaggac	660
cgggatg	ggca	caaagtgagt	gctcaccaaa	gcttgactgt	cctttcccat	ggcaatttac	720
ttcagct	tgt	ttgatttccc	ctccccgact	ggactaggca	cctattctct	gtcttctctc	780

tttacagttg gaaggagcaa aatgggactt ttggctgaaa gtgctgagct cctgcggtgg 840

gggctgacc	g caageegege	cttctgtgca	cctggtcggc	ccagctagct	gcggacccgg	900				
cggggaggg	id cadadcadac	caatcggcgc	tgccccagca	gggctgcggc	tgcaggcagc	960				
agagcctcc	t agecegtegg	tgtctgcgcc	catcgatccc	tttgtctatc	cccgaccatg	1020				
gcgaag						1026				
<210> 2 <211> 27										
<212> DN <213> Ho	omo sapiens									
<400> 2										
caaggaccg	gg gatggcacaa	agtgagtgct	caccaaagct	tgactgtcct	ttcccatggc	60				
aatttactt	c agcttgtttg	atttcccctc	cccgactgga	ctaggcacct	attctctgtc	120				
ttctctctt	t acagttggaa	ggagcaaaat	gggacttttg	gctgaaagtg	ctgagctcct	180				
gcggtgggg	gg ctgaccgcaa	gccgcgcctt	ctgtgcacct	ggtcggccca	gctagctgcg	240				
gacccggcg	ıd dd a ddddcdd	ggcgggccaa	tcgg			274				
<210> 3										
<211> 17										
<212> DN <213> Ar	IA tificial Sequ	ıence								
<220>										
<223> PC	DN1 SNP Prime	r 1								
<400> 3										
cegattggcc cgccccg 17										
<210> 4 <211> 17	,									
<212> DN										
<213> Ar	tificial Sequ	lence								
<220>										
<223> PC	N1 SNP Prime	r 2								
<400> 4										
ccgattggc	c cgcccca					17				
<210> 5										
<211> 18	3									
<212> DN	IA									
<213> Ar	tificial Seco	ience								

```
<223> PON1-107 consensus primer
<400> 5
caaggaccgg gatggcac
                                                                     18
<210> 6
<211> 19
<212> DNA
<213> Artificial Sequence
<220>
<223> Forward control primer coding for a fragment of the HLA-DRB3 gene
<400> 6
                                                                     19
tgccaagtgg agcacccaa
<210> 7
<211> 20
<212> DNA
<213> Artificial Sequence
<220>
<223> Reverse control primer coding for a fragment of the HLA-DRB3 gene
<400> 7
                                                                     20
gcatcttgct ctgtgcagat
<210> 8
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Reverse allele-specific primer for the PON1 M54L polymorphism
<400> 8
cagaaactgg ctctgaagac a
                                                                     21
<210> 9
<211> 21
<212> DNA
<213> Artificial Sequence
<220>
<223> Reverse allele-specific primer for the PON1 M54L polymorphism
<400> 9
cagaaactgg ctctgaagac t
                                                                     21
<210> 10
<211> 20
```

<212> DNA

<213> Artificial Sequence <220> <223> M54L Consensus primer

<400> 10

aagtgggcat gggtatacag

20